Supplemental Information

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Supplemental Data:

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- 5 Table S1, related to Figure 1. Characterization of the most potent D-enantiomeric peptides
- 6 against Pseudomonas aeruginosa PA14 biofilms. MBIC₅₀ values were determined using the 96-
- 7 well plate biofilm assay and correspond to the ability of the peptides to prevent biofilm formation
- 8 by 50%. The lowest MBIC₅₀ values obtained are shown in bold.

Peptide	Amino acid sequence	MIC (μg/ml)	MBIC ₅₀ (µg/ml)		
name					
DJK1	VFLRRIRVIVIR	20	10		
DJK2	VFWRRIRVWVIR	10	5		
DJK5	VQWRAIRVRVIR	16	1		
DJK6	VQWRRIRVWVIR	16	0.5		
RI-1002	KRIRWVILWRQV	10	5		
RI-1018	RRWIRVAVILRV	20	10		

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Species Strains Phenotype (μg/ml) (μg	concentration r	ange.		1010		DIVE		DIV	
Species		Do otomia i		1018		DJK5		DJK6	
ATTC 17987 Wild type >128 8	G		Diameter						
SENTRY C2 XDR; PXR >128 32 16 8 8 4			_	,, ,			(µg/mi)		(µg/mi)
SENTRY C5 MDR; PXR >128 128 16 8 16 4			V 1				1		1
SENTRY C8			· ·						
SENTRY C11 MDR; PXR >128 64 8 4 8 4 SENTRY C12 MDR; PXR >128 128 16 8 8 4 SENTRY C13 MDR; PXR >128 64 16 8 16 4 SENTRY C13 MDR; PXR >128 64 16 8 16 4 SENTRY C15 MDR; PXR >128 64 16 8 8 2 SENTRY C15 MDR; PXR >128 64 16 8 16 4 SENTRY C15 MDR; PXR >128 64 16 8 16 4 4 1 MDR; PXR >128 64 16 8 16 4 4 1 MDR; PXR >128 64 16 8 16 4 4 1 MDR; PXR >128 4 8 4 4 1 MDR MDR >256 2 >256 0.4 >64 2 2 2 2 2 2 2 2 2									
SENTRY C12 MDR; PXR >128 128 16 8 8 4			, , , , , , , , , , , , , , , , , , ,						
SENTRY C13 MDR; PXR >128 64 16 8 8 16 4 SENTRY C14 DR; PXR >128 16 16 8 8 8 2 SENTRY C15 MDR; PXR >128 64 16 8 16 4 SENTRY C87 XDR; PXR >128 4 8 4 4 1 Burkholderia 4813 MDR >256 2 >256 0.4 >64 2 Enterobacter cloacae 218R Derepressed 64 2 32 2 16 2 Escherichia Coli HB101 WT 32 8 1.6 0.8 16 8 Escherichia Coli HB101 WT 32 16 16 4 8 2 VGH10 MDR 256 1 8 4 8 8 VGH1 MDR 64 32 16 4 8 8 VGH1 MDR 64 32 8 2 16 8 Klebsiella KPLN649 WT 32 1 3.2 1.6 4 2 Pseudomonas PA01 WT 64 8 16 2 16 1 Brazil #21 MDR; PXR 64 8 16 4 >16 16 Brazil #36 MDR; PXR 64 8 16 4 >16 16 Brazil #212 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #212 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #213 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #214 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #215 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #210 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #211 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #211 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #211 MDR; PXR 64 32 >16 0.5 8 0.5 LES400 - 27 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1		SENTRY C11	MDR; PXR	>128	64	8			4
SENTRY C14 DR; PXR >128 16 16 8 8 16 4		SENTRY C12	MDR; PXR	>128	128	16	8	8	4
SENTRY C15 MDR; PXR >128 64 16 8 16 4		SENTRY C13	MDR; PXR	>128	64	16	8	16	4
SENTRY C87 XDR; PXR >128 4 8 4 4 1		SENTRY C14	DR; PXR	>128	16	16	8	8	2
Burkholderia cenocepacia		SENTRY C15	MDR; PXR	>128	64	16	8	16	4
cenocepacia 218R Derepressed Class C β-lactamase 64 2 32 2 16 2 Escherichia coli 0157 WT 32 8 1.6 0.8 16 8 HB101 WT 32 16 16 4 8 2 VGH10 MDR 256 1 8 4 8 2 VGH11 MDR 64 32 16 4 8 8 VGH1 MDR 64 32 8 2 16 8 Klebsiella pneumoniae KPLN649 WT 32 1 3.2 1.6 4 2 Pseudomonas aeruginosa PA01 WT 64 8 >64 2 32 2 Pseudomonas aeruginosa PA14 WT 32 8 16 1 16 0.5 Brazil #21 MDR; PXR 64 8 16 1 16 0.5 B		SENTRY C87	XDR; PXR	>128	4	8	4	4	1
Enterobacter cloacae	Burkholderia	4813	MDR	>256	2	>256	0.4	>64	2
cloacae Class C β-lactamase Class C β-lactamase Escherichia coli 0157 WT 32 8 1.6 0.8 16 8 WGH10 WT 32 16 16 4 8 2 VGH10 MDR 256 1 8 4 8 2 VGH11 MDR 64 32 16 4 8 8 VGH1 MDR 64 32 8 2 16 8 Klebsiella pneumoniae KPLN649 WT 32 1 3.2 1.6 4 2 Pseudomonas aeruginosa PA01 WT 64 8 16 2 16 1 PA14 WT 32 8 16 1 16 0.5 Brazil #21 MDR; PXR 64 8 16 4 >16 1 Brazil #36 MDR; PXR 64 4 8 0.5 16 1	cenocepacia								
Escherichia Coli HB101 WT 32 8 1.6 0.8 16 8 16 WT 32 16 16 4 8 2 VGH10 MDR 256 1 8 4 8 2 VGH11 MDR 64 32 16 4 8 8 8 VGH1 MDR 64 32 8 2 16 8 8 8 VGH1 MDR 64 32 8 2 16 8 8 8 VGH1 MDR 64 32 8 2 16 8 8 8 8 8 8 8 8 8	Enterobacter	218R	Derepressed	64	2	32	2	16	2
Bescherichia Coli HB101 WT 32 8 1.6 0.8 16 8 1.6 0.8 16 8 1.6 0.8 1.6 1.	cloacae		Class C β-						
HB101									
VGH10	Escherichia	0157	WT	32	8	1.6	0.8	16	8
VGH11 MDR 64 32 16 4 8 8 VGH1 MDR 64 32 8 2 16 8 Klebsiella pneumoniae KPLN649 WT 32 1 3.2 1.6 4 2 Pseudomonas aeruginosa PA01 WT 64 8 16 2 32 2 Pseudomonas aeruginosa PA14 WT 32 8 16 1 16 0.5 Brazil #21 MDR; PXR 64 8 16 1 16 0.5 Brazil #36 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #172 MDR; PXR 64 4 8 0.5 16 1 Brazil #205 MDR; PXR 64 4 8 0.5 16 8 Brazil #211 MDR; PXR 64 4 8 0.5 16 8 Brazil #200 30	coli	HB101	WT	32	16	16	4	8	2
VGH1 MDR 64 32 8 2 16 8 Klebsiella KPLN649 WT 32 1 3.2 1.6 4 2 pneumoniae 1549216 WT >64 8 >64 2 32 2 Pseudomonas PA01 WT 64 8 16 2 16 1 aeruginosa PA14 WT 32 8 16 1 16 0.5 Brazil #21 MDR; PXR 64 8 16 4 >16 16 Brazil #36 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #172 MDR; PXR 64 4 8 0.5 16 1 Brazil #205 MDR; PXR 64 4 8 0.5 16 1 Brazil #211 MDR; PXR 64 4 8 0.5 16 8 Brazil #211 MDR; PXR 64 32 >16 0.5 8 0.5 LES400 - 27 MDR 16 2 8 2 4 1 LES400 - 30 MDR 16 1 8 1 4 0.5 LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1		VGH10	MDR	256	1	8	4	8	2
Klebsiella		VGH11	MDR	64	32	16	4	8	8
Decimoniae 1549216 WT >64 8 >64 2 32 2		VGH1	MDR	64	32	8	2	16	8
Pseudomonas aeruginosa PA01 WT 64 8 16 2 16 1 Brazil #36 MDR; PXR 64 8 16 1 16 0.5 Brazil #36 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #36 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #172 MDR; PXR 64 32 >16 0.5 16 1 Brazil #205 MDR; PXR 64 4 8 0.5 16 8 Brazil #211 MDR; PXR 64 32 >16 0.5 8 0.5 LES400 - 27 MDR 16 2 8 2 4 1 LES400 - 30 MDR 16 1 8 1 4 0.5 LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64	Klebsiella	KPLN649	WT	32	1	3.2	1.6	4	2
PA14 WT 32 8 16 1 16 0.5 Brazil #21 MDR; PXR 64 8 16 4 >16 16 Brazil #36 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #172 MDR; PXR 64 32 >16 0.5 16 1 Brazil #205 MDR; PXR 64 4 8 0.5 16 8 Brazil #211 MDR; PXR 64 32 >16 0.5 8 0.5 LES400 - 27 MDR 16 2 8 2 4 1 LES400 - 30 MDR 16 1 8 1 4 0.5 LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1	pneumoniae	1549216	WT	>64	8	>64	2	32	2
Brazil #21 MDR; PXR 64 8 16 4 >16 16 Brazil #36 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #172 MDR; PXR 64 32 >16 0.5 16 1 Brazil #205 MDR; PXR 64 4 8 0.5 16 8 Brazil #211 MDR; PXR 64 32 >16 0.5 8 0.5 LES400 - 27 MDR 16 2 8 2 4 1 LES400 - 30 MDR 16 1 8 1 4 0.5 LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1	Pseudomonas	PA01	WT	64	8	16	2	16	1
Brazil #36 MDR; PXR 64 4 8 0.5 16 0.5 Brazil #172 MDR; PXR 64 32 >16 0.5 16 1 Brazil #205 MDR; PXR 64 4 8 0.5 16 8 Brazil #211 MDR; PXR 64 32 >16 0.5 8 0.5 LES400 - 27 MDR 16 2 8 2 4 1 LES400 - 30 MDR 16 1 8 1 4 0.5 LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1	aeruginosa	PA14	WT	32	8	16	1	16	0.5
Brazil #172 MDR; PXR 64 32 >16 0.5 16 1 Brazil #205 MDR; PXR 64 4 8 0.5 16 8 Brazil #211 MDR; PXR 64 32 >16 0.5 8 0.5 LES400 - 27 MDR 16 2 8 2 4 1 LES400 - 30 MDR 16 1 8 1 4 0.5 LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1		Brazil #21	MDR; PXR	64	8	16	4	>16	16
Brazil #205 MDR; PXR 64 4 8 0.5 16 8 Brazil #211 MDR; PXR 64 32 >16 0.5 8 0.5 LES400 - 27 MDR 16 2 8 2 4 1 LES400 - 30 MDR 16 1 8 1 4 0.5 LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1		Brazil #36	MDR; PXR	64	4	8	0.5	16	0.5
Brazil #211 MDR; PXR 64 32 >16 0.5 8 0.5 LES400 - 27 MDR 16 2 8 2 4 1 LES400 - 30 MDR 16 1 8 1 4 0.5 LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1		Brazil #172	MDR; PXR	64	32	>16	0.5	16	1
LES400 - 27 MDR 16 2 8 2 4 1 LES400 - 30 MDR 16 1 8 1 4 0.5 LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1		Brazil #205	MDR; PXR	64	4	8	0.5	16	8
LES400 - 30 MDR 16 1 8 1 4 0.5 LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1		Brazil #211	MDR; PXR	64	32	>16	0.5	8	0.5
LES400 - 31 MDR 16 2 4 1 4 2 Salmonella C587 WT 64 3.2 3.2 0.8 4 1			MDR	16	2	8	2	4	1
Salmonella C587 WT 64 3.2 3.2 0.8 4 1		LES400 - 30	MDR	16	1	8	1	4	0.5
Salmonella C587 WT 64 3.2 3.2 0.8 4 1		LES400 - 31	MDR	16	2	4	1	4	2
enterica	Salmonella	C587	WT	64	3.2	3.2	0.8	4	1
	enterica								

Figure S1, related to Figure 4. Overproduction of (p)ppGpp led to decreased biofilm susceptibility to D-enantiomeric peptides. Addition of SHX, which leads to overproduction of (p)ppGpp, resulted in the resistance of flow cell biofilm formation to 2.5 μg/ml of peptide DJK-5. After 3 days, bacteria were stained green with the all bacteria stain Syto-9 prior to confocal imaging. Each panel shows reconstructions from the top in the large panel and sides in the right and bottom panels (xy, yz and xz dimensions).

